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	10	20	30	40	50	60	
1	HHNGTNGTMMQYFEWYLPNDGNHWNRLRDDAANLKS KGITAVWI PPAWKGSQNDVG YGA						60
3	-AAPFNGTMMQYFEWYLPDDGTLWTKVANEANNLSSLGITALWLPPAYKGT SRSDVG YGV						59
2	HHNGTNGTMMQYFEWHL PNDGNHWNRLRDDASNLRNRGITAIWI PPAWKGSQNDVG YGA						60
4	HHNGTNGTMMQYFEWYLPNDGNHWNRLNSDASNLSKSGITAVWI PPAWK GASQNDVG YGA						60
	70	80	90	100	110	120	
1	YDLYDLGEFNQKGTVRTKYGTRNQLQAAVTS LKNNGIQVYGDVVMNHKGGADGTEI VNAV						120
3	YDLYDLGEFNQKGTVRTKYGTAKAYLQAIQAAHAAGMQVYADVVF DHKGGADGTEWV DAV						119
2	YDLYDLGEFNQKGTVRTKYGTRS QLES AIHALKNGVQVYGDVVMNHKGGADATEN V LAV						120
4	YDLYDLGEFNQKGTVRTKYGTRS QLQAAVTS LKNNGIQVYGDVVMNHKGGADATE M VRAV						120
	130	140	150	160	170	180	
1	EVNRSNRNQETS GEYAI EAWTKFDFPGRGNHSSFKWRWYHFDGTDWDQSRQLQNKIYKF						180
3	EVNPSDRNQEIS GTYQIQA WTKFDFPGRGNTYSSFKWRWYHFDGVDWDESRKLS -RIYKF						178
2	EVNPNNRNQEIS GDTIEAWTKFDFPGRGNTYSDFKWRWYHFDGVDWDQSRQFQNR IYKF						180
4	EVNPNNRNQEV TGEYTI EAWTRFDFPGRGNTHSSFKWRWYHFDGVDWDQSRRLN NRIYKF						180
	190	200	210	220	230	240	
1	RGTGKAWDWEVD TENGNYDYL MYADVDMDHPEVIHEL RNWGVWYTTNTLNL DGFRI DAVKH						240
3	RGIGKAWDWEVD TENGNYDYL MYADLMDHPEVVT ELKNWGK WYVNTTNIDGFR L DAVKH						238
2	RGDGKAWDWEVDSENGNYDYL MYADVDMDHPEV VNELRRWGEWYTTNTLNL DGFRI DAVKH						240
4	RGHGKAWDWEVD TENGNYDYL MYADIDMDHPEV VNELRNWGVWYTTNTLGL DGFRI DAVKH						240
	250	260	270	280	290	300	
1	IKYSFTRDWLTHVRNTTGKPMFAVAEFWKNDLGAIENYLNKTSWNHSAFDVPLHYNLYNA						300
3	IKFSFFPDWLSYVRSQTGKPLFTVGEYWSYDINKLHNYITKTDGTM SLFDAPLHNKFYTA						298
2	IKYSFTRDWLTHVRNATGKEMFAVAEFWKNDLGAIENYLNKTNWNH SVFDVPLHYNLYNA						300
4	IKYSFTRDWINHVR SATGKNMFAVAEFWKNDLGAIENYLQKTNNH SVFDVPLHYNLYNA						300
	310	320	330	340	350	360	
1	SNSGGYDMRNILNGSVVQKHPH AVTFVDNHDSQPGEALESFVQQWF KPLAYALV LTRI						360
3	SKSGGAFDMRTLMTNTL MKDQPTLAVTFVDNH DTEPGQALQSWVDPWF KPLAYA FILTRQ						358
2	SNSGGNYDMAKLLNGTVVQKHPMHA VTFVDNHDSQPGELESFVQEWF KPLAYAL ILTRE						360
4	SKSGGNYDMRNIFNGTVVQRHP SHAVTFVDNHDSQPEEAL ESFVEEWF KPLAYAL TLTRE						360
	370	380	390	400	410	420	
1	QGYPVS VFYGDYYGIP THGVPAMKSKIDP LLQARQTFAYGTQHDYFDHDI IGWTREGNSS						420
3	EGYPCVFYGDYYGIPQYNI PSLKSKIDP LLIARRDYAYGTQHDYLDHSDI IGWTREGGTE						418
2	QGYPVS VFYGDYYGIP THSVPAMKAKIDP ILEARQNFAYGTQHDYFDHNI IGWTREGNTT						420
4	QGYPVS VFYGDYYGIP THGVPAMRSKIDP ILEARQKYAYGKQNDYLDHNI IGWTREGNTA						420
	430	440	450	460	470	480	
1	HPNSGLATIMSDGPGGNKWMYVGK NKAGQVWRDITGNRTGTVTINADGWGNF SVNGGSVS						480
3	KPGSGLAALITDGPGGSKWMYVGK QHAGKVFYDLTGNRSDTVTINSDGWGEFKVNGGSVS						478
2	HPNSGLATIMSDGPGGEKWMYVGQNKAGQVWSDITGNKPGTVTINADGWANF SVNGGSVS						480
4	HPNSGLATIMSDGAGGSKWMFVGRNKAGQVWSDITGNRTGTVTINADGWGNF SVNGGSVS						480
	490	500	510	520	530	540	
1	VWVKQ						485
3	VWVPRKTTVSTIARPITTRPWTGEFVRWTEPRLVAV						514
2	IWKVR						485
4	IWNK						485

Fig. 1

09902188-071001

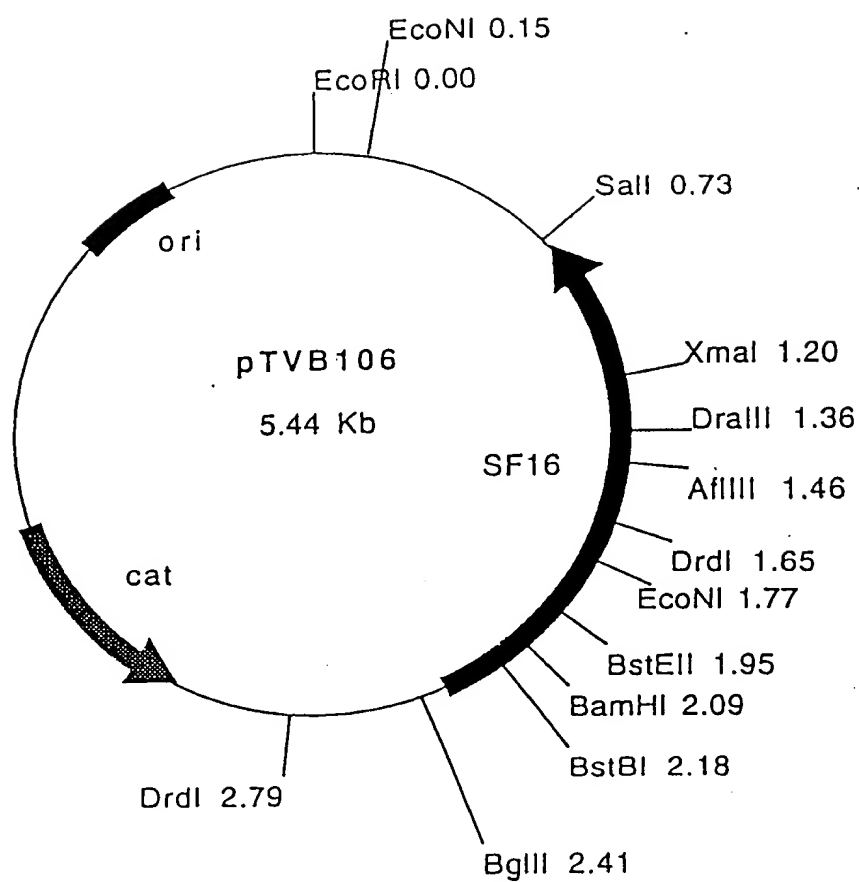


Fig. 2

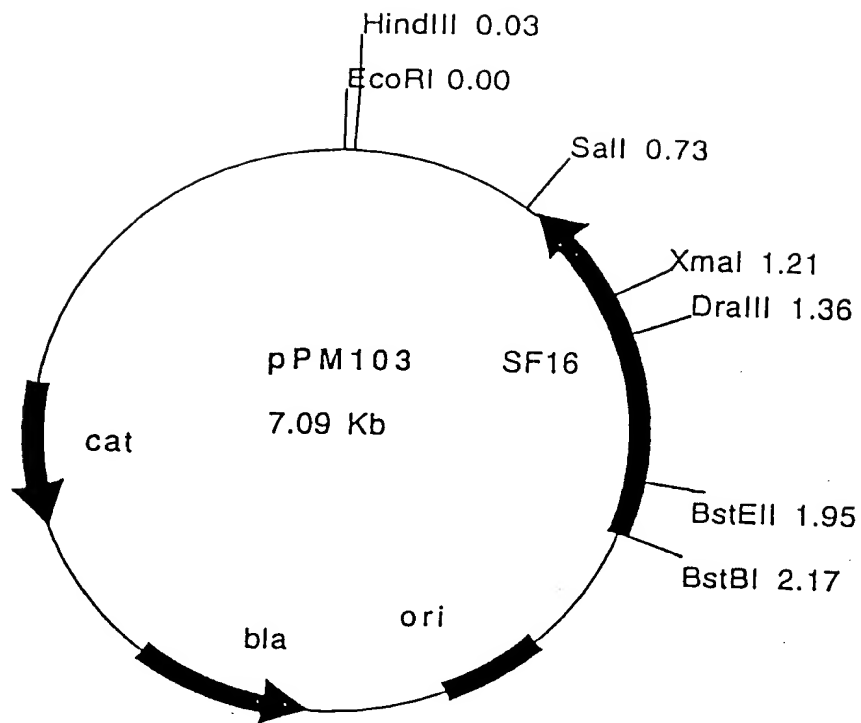


Fig. 3

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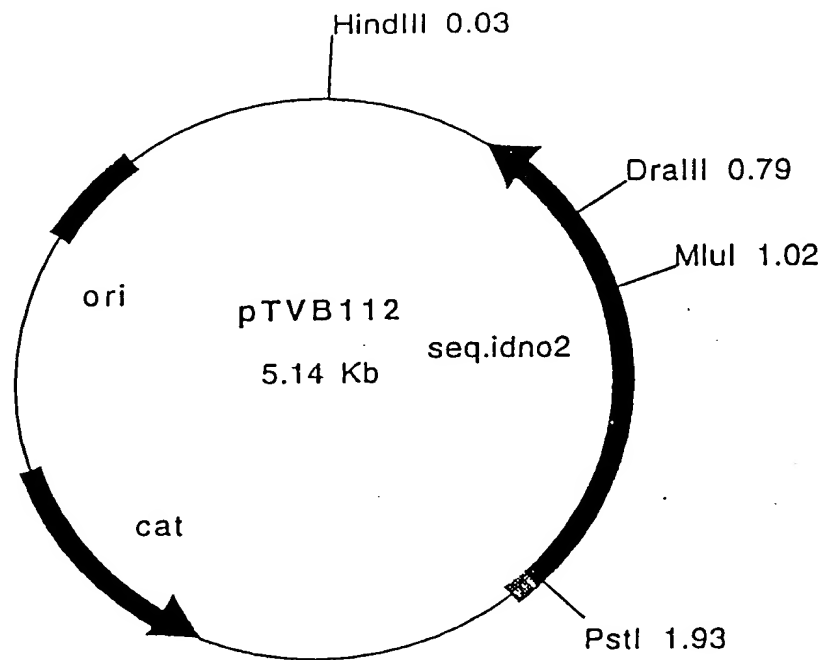


Fig. 4

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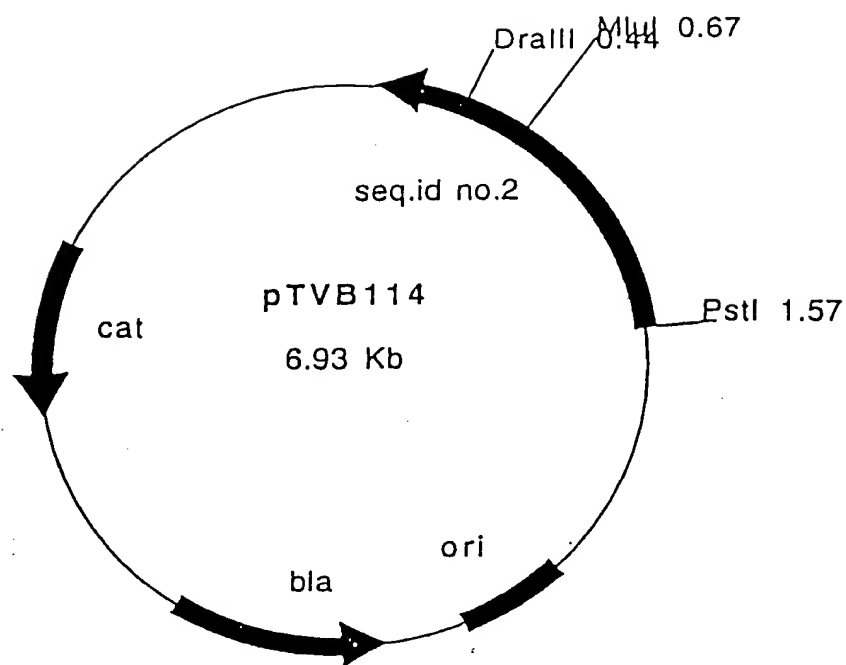


Fig. 5

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